

Fig. 1A

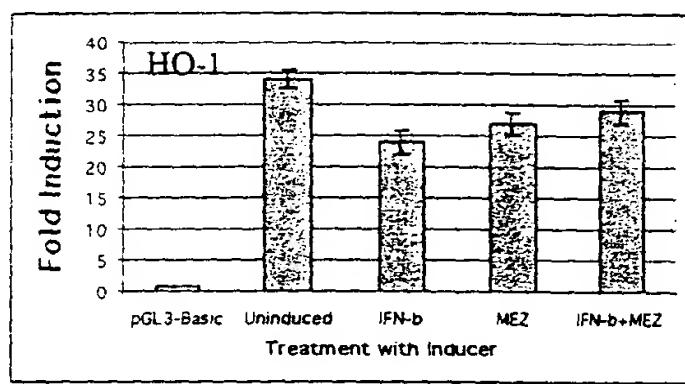


Fig. 1B

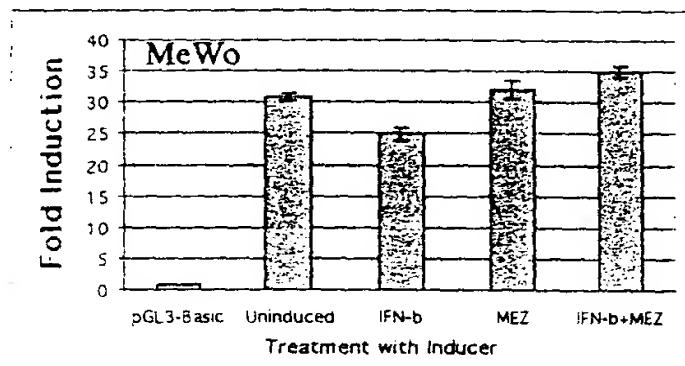


Fig. 2A**Fig. 2B****Fig. 2C**

Hu. mda-7 UUGUAUUAUACAACUCUAUUAAUGUCAGUAUUCAACUGAACGUUUUUU
 Hu. α -IFN UAUUUUUUUUUUAA
 Hu. GM-CSF UAUUUUUUUUUUUUUUUAA
 Hu. TNF AUUUUUUUUUUUUUUUAA
 Hu. cFos GUUUUUUUUUUUUUUUUUAAAGAUGGAUUCAGAUUUUUUUUUUUUUUU

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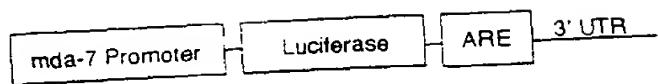


Fig. 3A

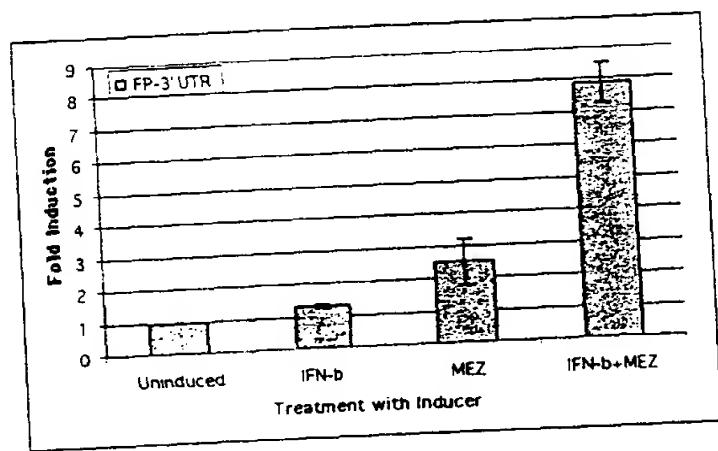
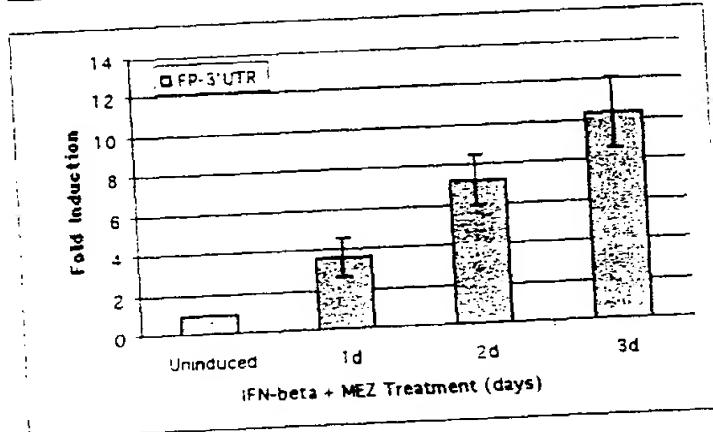
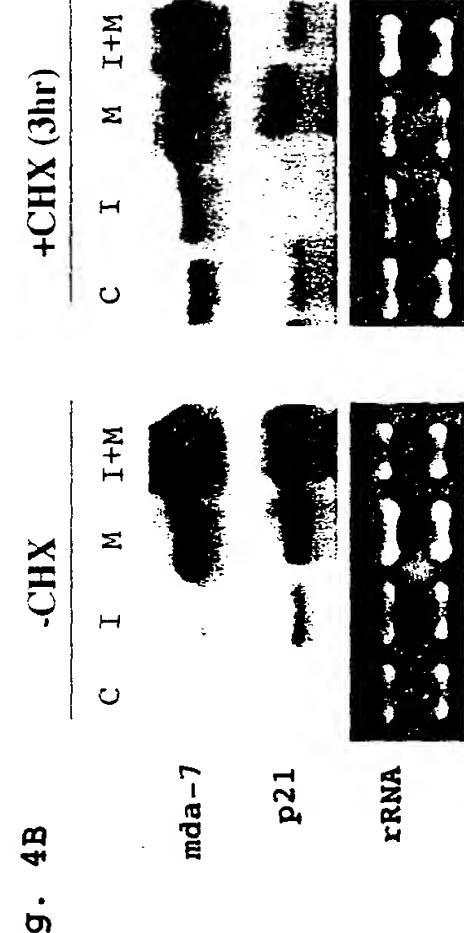
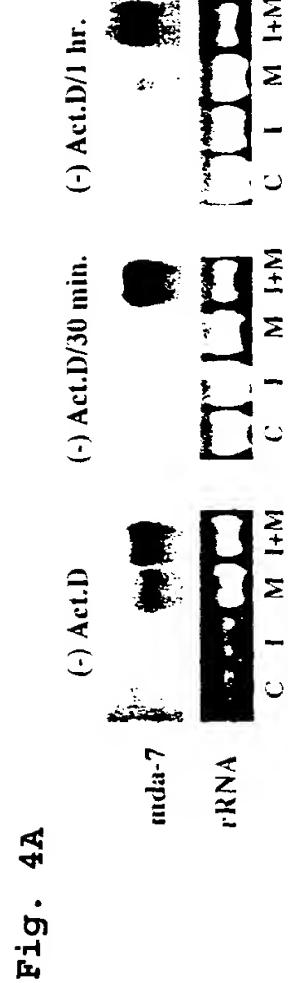


Fig. 3B





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Fig. 5A

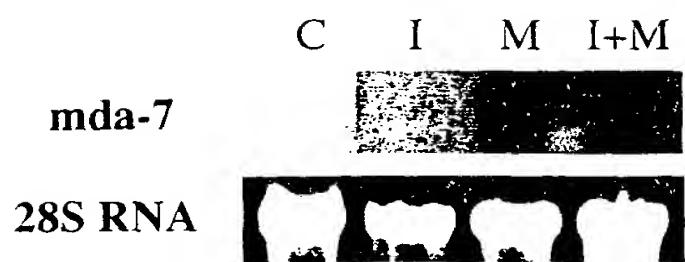


Fig. 5B

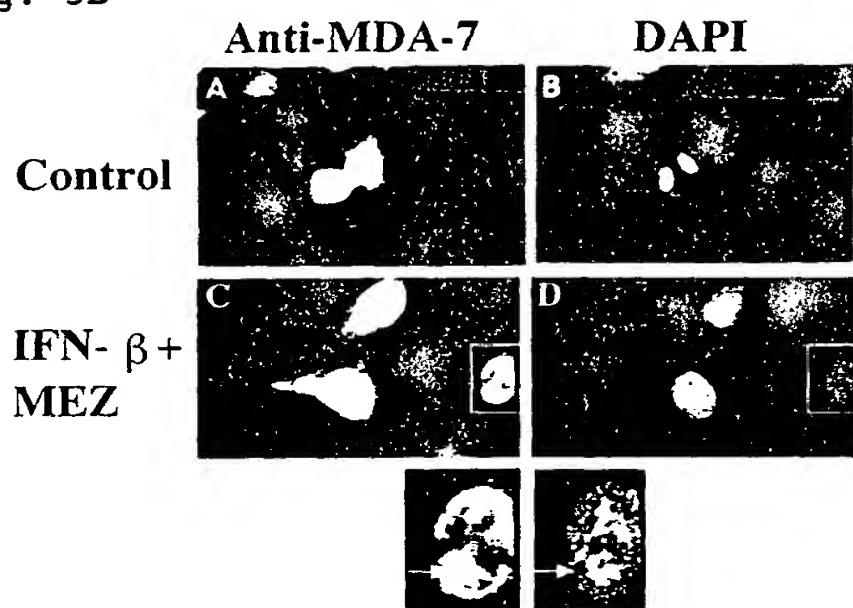


Fig. 6

TAATACGACTCACTATAGGGCGTCGACTCGATCACCTTTGAACCCAGGTCTGCCTGCCTCAAAGCTTGTACTCATAAC - 2161
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 C/EBP
 CTCATCGGAGGCTGAGGCCAGGCACATGTTGCCTGAACATCCATGTTATATGATTCCCTCAGACAGAGTGAGCT - 2001
 ACTCACGATCCCAGGTGTACCCCTGAGGCCAGCAAGGTGTATCCATGACCTCATGCCTCTGTTCCAGCCTGCCCTTAAC - 1921
 AGCTCATCCCACCTGCCTGCCCTCCCCCTATCTGCAGACAGTAGTCTAGGATTCAAGCTGCCCTGGGGCTCATTTC - 1841
 CCTCTCAGCTTCTGCTTAGCTGTCTGCCTCCACTCACCTATTACTCCAGCACTCTCACCTGGTCTTCTG - 1761
 C/EBP
 TCTCATCACTGCCTCTTGACATCTTATCTCATAGTAGTTAGGGTTCTGGTAATGCCCTAAATCCACATGGTGG - 1681
 GAAGGGGGAGTGGGGAAAGAGAGTGCCTGTGGCTGTGCCTACTCTGGAGGGTAAGACTCGGGCCCTCAGGAACA - 1601
 AAGGATTCAAGCTGGTGGCAGCTATGCCAAGCAGACTGCTGCCAGGGATTGCAAAGGAGTATTTGTTGCTTAAGAA - 1521
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 C/EBP
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 AP1
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 C/EBP
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 AP1 C/EBP
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 AP1
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 GTGTTGTTTGCCTTGTATTACACTGGGACAAATTAAAATTATACATGCAGAGACTGCAGCGCAGAGAAC - 241
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 AP1 C/EBP
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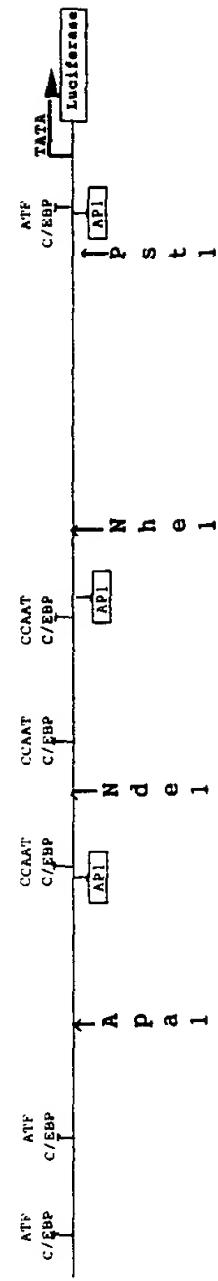
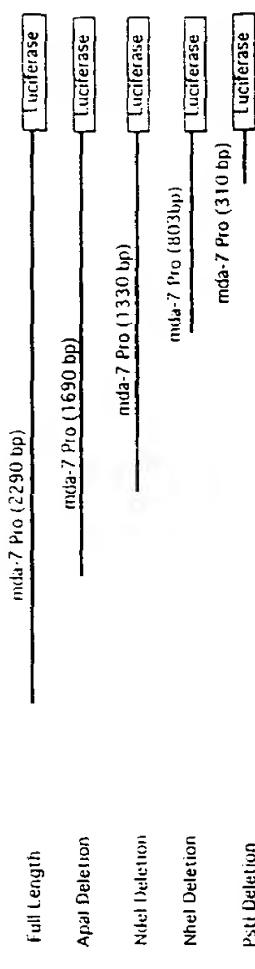
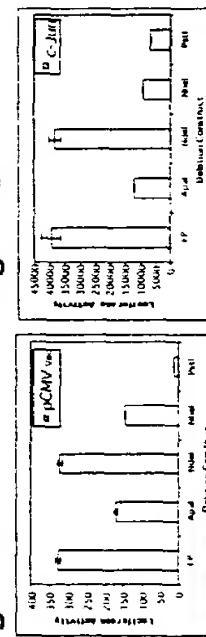
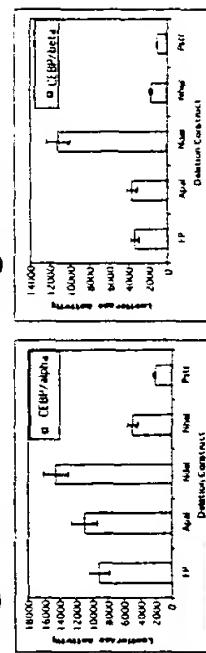
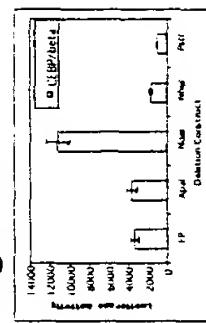
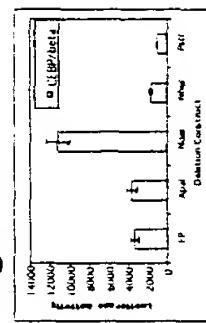
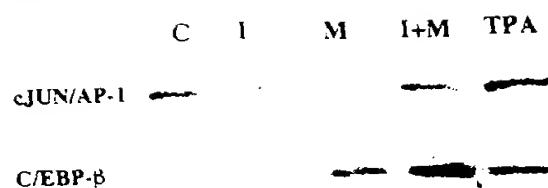
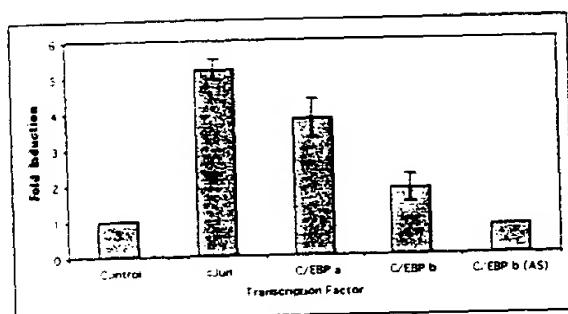
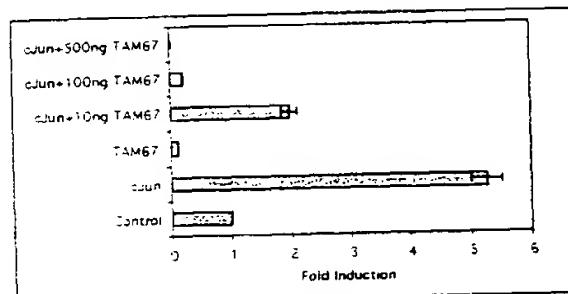
Fig. 7A**Fig. 7B****Fig. 7C****Fig. 7D****Fig. 7E****Fig. 7F**

Fig. 8A**Fig. 8B****Fig. 8C**

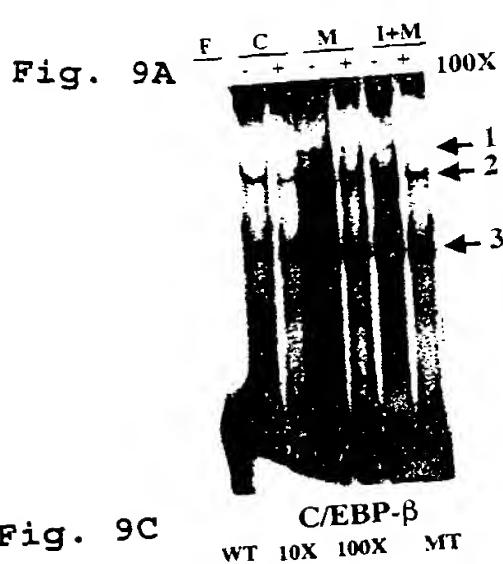


Fig. 9C

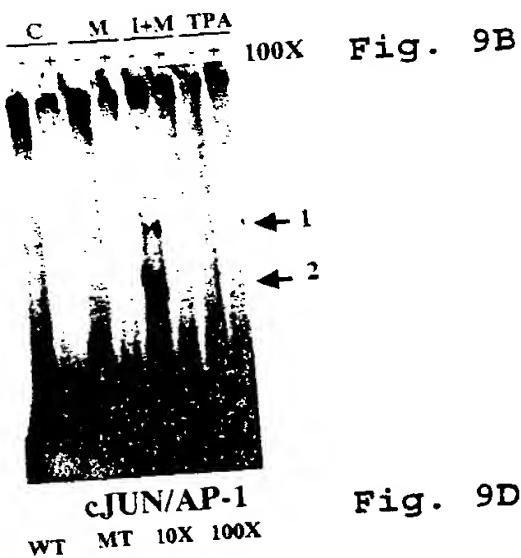


Fig. 9D



Fig. 9E

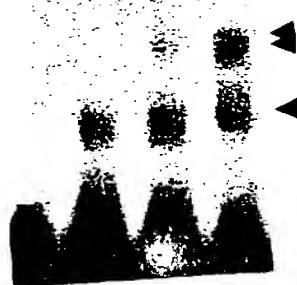


Fig. 9F

